

# **Combining Information: Heteroscedastic Random-Effects Models for Interlaboratory Comparisons**

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## **Interlaboratory Studies: The Scenario**

- Each of  $p$  laboratories makes repeated measurements of  $m$  quantities (perhaps corresponding to different concentrations of a chemical analyte).
- The number of measurements made can differ among the laboratories.
- The measurement variability may depend on the material being measured (perhaps as an increasing function of concentration or level).
- The within-laboratory variabilities may differ (often, though, they are assumed to be equal).

## **Interlaboratory Studies: Some questions**

- How should one estimate 'consensus' values of the quantities measured?
- What is the between-laboratory variability (*reproducibility*)?
- What is the within-laboratory variability (*repeatability*)? How do they compare?
- How should we look for outliers?

## **Why Interlaboratory Studies?**

- Interlaboratory studies are primarily performed for one of two reasons:
  1. Validating a measurement method or standard material
  2. Assessing the proficiency of measurement laboratories.

## Outline

- A single material measured by multiple laboratories – *one-way random model* (heteroscedastic and unbalanced)
  - Likelihood Analysis
  - Bayesian Model and Credible Regions
  - Example
- Some results for two-way models.

## Dietary Fiber in Apricots Li and Cardozo (1994)

Lab.	$x_i$	$s_i^2$	$n_i$
1	25.32	0.37	2
2	26.72	0.62	2
3	27.89	0.35	2
4	27.70	1.85	2
5	27.42	0.61	2
6	24.30	0.21	2
7	27.11	0.37	2
8	27.28	0.09	2
9	25.37	0.08	2

Mean:  $\bar{x} = 26.567$

Weighted Means:

MP    =    26.472  
 GD    =    26.164  
 ANOVA =    26.420  
 MLE   =    27.275

## Statistical Framework: One-Way, Unbalanced, Heteroscedastic Random-Effects ANOVA

- Laboratory sample means  $x_i$  distributed independently normal with mean  $\mu$  and variance  $\sigma^2 + \tau_i^2$ , where  $\tau_i^2 = \sigma_i^2/n_i$ .
- Expected mean for  $i$ th laboratory is also normal, with mean  $\mu$  and variance  $\sigma^2$ .
- Sufficient statistics  $x_i$  and  $t_i^2 = s_i^2/n_i$ .

If  $x_{ij}$  denotes the  $j$ th measurement from the  $i$ th lab, then

$$x_{ij} = \mu + b_i + e_{ij},$$

where  $b_i \sim N(0, \sigma^2)$  and  $e_{ij} \sim N(0, \sigma_i^2)$ ; mutually independent.

## Maximum Likelihood (Cochran, 1937)

Let  $\omega_i = 1/(\sigma^2 + \tau_i^2)$ ,  $\nu_i = n_i - 1$ , and determine  $\hat{\sigma}$ ,  $\hat{\tau}_i^2$ , and  $\hat{\mu}$  to satisfy

$$(A_i) \quad \omega_i - \omega_i^2(x_i - \mu)^2 + \nu_i \left( \frac{1}{\tau_i^2} - \frac{t_i^2}{\tau_i^4} \right) = 0$$

$$(B) \quad \boxed{\sum_{i=1}^k \omega_i^2(x_i - \mu)^2 = \sum_{i=1}^k \omega_i}$$

$$(C) \quad \mu = \frac{\sum_{i=1}^k \omega_i x_i}{\sum_{i=1}^k \omega_i}$$

Note that (B) may have multiple roots. Cochran (1937) proposed setting  $\tau_i^2 = t_i^2$  and solving (B) for  $\sigma^2$ , then using (C).

## ML Equations

$$\mu = \frac{\sum_{i=1}^p \gamma_i x_i}{\sum_i \gamma_i} = \frac{\sum_{i=1}^p \omega_i x_i}{\sum_i \omega_i}$$

$$\sigma^2 = \frac{\sum_{i=1}^p \gamma_i \left[ (x_i - \mu)^2 + \frac{\nu_i t_i^2}{1 - \gamma_i} \right]}{\sum_{i=1}^p n_i}$$

$$\begin{aligned} &\gamma_i^3 - (a_i + 2)\gamma_i^2 + \\ &[(n_i + 1)a_i + (n_i - 1)b_i + 1] \gamma_i \\ &- n_i a_i = 0 \end{aligned}$$

where

$$\gamma_i \equiv \frac{\sigma^2}{\sigma^2 + \tau_i^2}$$

$$a_i \equiv \frac{\sigma^2}{(x_i - \mu)^2}$$

and

$$b_i \equiv \frac{t_i^2}{(x_i - \mu)^2}.$$

## Result #1: Monotone Convergence to Stationary Points of the Likelihood

- For any starting values  $\mu_0, \sigma_0^2$ , maximize the likelihood over the weights by solving the cubics. (If there are multiple real roots, choose the one which causes the biggest increase in the likelihood.)
- Let

$$\sigma_1^2 = \frac{\sum_{i=1}^p \gamma_i \left[ (x_i - \mu)^2 + \frac{\nu_i t_i^2}{1 - \gamma_i} \right]}{\sum_{i=1}^p n_i}$$
$$\mu_1 = \frac{\sum_{i=1}^p \gamma_i x_i}{\sum_{i=1}^p \gamma_i}$$

solve for new weights, and iterate.

- This iteration, *regardless of starting values*, always converges to a stationary point of the likelihood, and *increases the likelihood at each step*.

## Result #2: Location of Stationary Values of the Likelihood

- At a stationary point of the likelihood,

$$\hat{\sigma}^2 = \frac{\sum_{i=1}^p \gamma_i^2 (x_i - \mu)^2}{\sum_{i=1}^p \gamma_i}$$

hence

- *All* of the stationary points of the likelihood  $\hat{\mu}$  and  $\hat{\sigma}$  are within the rectangle in the  $(\mu, \sigma)$  plane given by

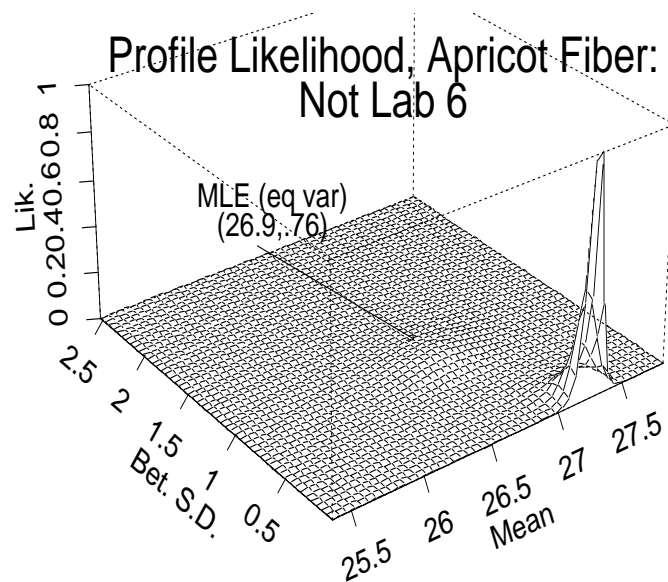
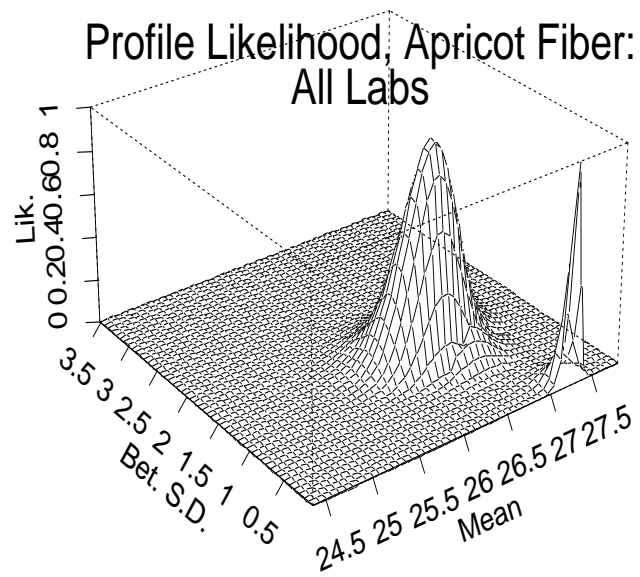
$$\min_i(x_i) \leq \tilde{\mu} \leq \max_i(x_i)$$

and

$$0 \leq \tilde{\sigma} \leq \max_i(x_i) - \min_i(x_i).$$

- After the appropriate location-scale transformation of the data, it is only necessary to search the unit square in the  $(\mu, \sigma)$  plane for stationary values.

## Lab. 6 an Outlier for Apricot Data



**Result #3:**  
**Location of the Roots of Cubic**  
**Equations for Weights ( $\gamma_i$ )**

- Each cubic likelihood equation has one or three roots  $\gamma_i \in [0, 1]$ .
- A necessary condition for three roots is that

$$(x_i - \mu)^2 \geq \max(\sigma^2/q_i, t_i^2/h_i),$$

where

$$\begin{aligned} q_i &= -2 - 6\sqrt{n_i} \sin \left\{ \frac{1}{3} \left[ \sin^{-1} \left( \sqrt{\frac{n_i - 1}{n_i}} \right) - \frac{\pi}{2} \right] \right\} \\ &= \frac{8}{27n_i} + O(n_i^{-2}) \end{aligned}$$

and

$$h_i = \frac{(1 - q_i)^3}{27(n_i - 1)} = \frac{1}{27n_i} + O(n_i^{-2}).$$

- These values  $q_i$  and  $h_i$  are the smallest for which this is necessary.

**One-Way Models in  
Interlaboratory Studies:  
The Mandel-Paule Estimator  
J. of Research of the NBS (1982)**

- For arbitrary positive weights  $\{w_i\}_{i=1}^k$ , weighted mean is

$$\tilde{\mu} = \frac{\sum_{i=1}^p w_i x_i}{\sum_{i=1}^p w_i}.$$

- *Mandel-Paule* estimate,  $\mu_{MP}$ , of  $\mu$  is the weighted mean  $\tilde{\mu}$  for which

$$w_i \equiv \frac{1}{\tilde{\sigma}^2 + t_i^2}$$

where  $\tilde{\sigma}^2$  is the root (if any) of

$$Q = \sum_{i=1}^p w_i (x_i - \tilde{\mu})^2 = p - 1$$

- Note:  $Q$  is convex decreasing on  $[0, \infty)$ , and  $Q \sim \chi_{p-1}^2$  if

$$w_i = \omega_i \equiv \frac{1}{\sigma^2 + \tau_i^2}$$

## The Mandel-Paule Algorithm and ML/REML

Maximum-Likelihood for a linear model

$$Y = X\beta + e,$$

where  $e \sim N(0, \Sigma)$  is equivalent to minimizing  $|\Sigma|$ , subject to

$$(y - X\hat{\beta})^T \Sigma^{-1} (y - X\hat{\beta}) = n \quad (1)$$

where  $\hat{\beta}$  is the GLS estimate of  $\beta$ , and  $n$  is the number of observations.

For our one-way model, if the  $\sigma_i^2$  are replaced by  $s_i^2$ , then (1), an equation in  $\sigma^2$  alone, is

$$\sum_{i=1}^p w_i (x_i - \tilde{\mu})^2 = p.$$

Had REML been used, rather than ML, then the  $p$  on the RHS above would be a  $p - 1$ , *precisely* Mandel and Paule's equation.

## Hierarchical Model With Noninformative Priors

$i = 1, \dots, p$  indexes laboratories

$j = 1, \dots, n_i$  indexes measurements

$$p(x_{ij}|\delta_i, \sigma_i^2) = N(\delta_i, \sigma_i^2)$$

$$p(\sigma_i) \propto 1/\sigma_i$$

$$p(\delta_i|\mu, \sigma^2) = N(\mu, \sigma^2)$$

$$p(\mu) = 1$$

$$p(\sigma) = 1$$

## A Useful Probability Density

Let  $T_\nu$  and  $Z$  denote independent Student- $t$  and standard normal random variables, and assume that  $\psi \geq 0$  and  $\nu > 0$ . Then

$$U = T_\nu + Z\sqrt{\frac{\psi}{2}}$$

has density

$$f_\nu(u; \psi) \equiv \frac{1}{\nu/2\sqrt{\pi}} \int_0^\infty \frac{y^{(\nu+1)/2-1} e^{-y\left[1+\frac{u^2}{\psi y+\nu}\right]}}{\sqrt{\psi y+\nu}} dy.$$

## Posterior of $(\mu, \sigma)$

- Assume  $\delta_i \sim N(\mu, \sigma^2)$ ,  $\sigma \sim p(\sigma)$ ,  
 $p(\mu) = 1$ ,  $p(\sigma_i) = 1/\sigma_i$ .

- Then the posterior of  $(\mu, \sigma)$  is

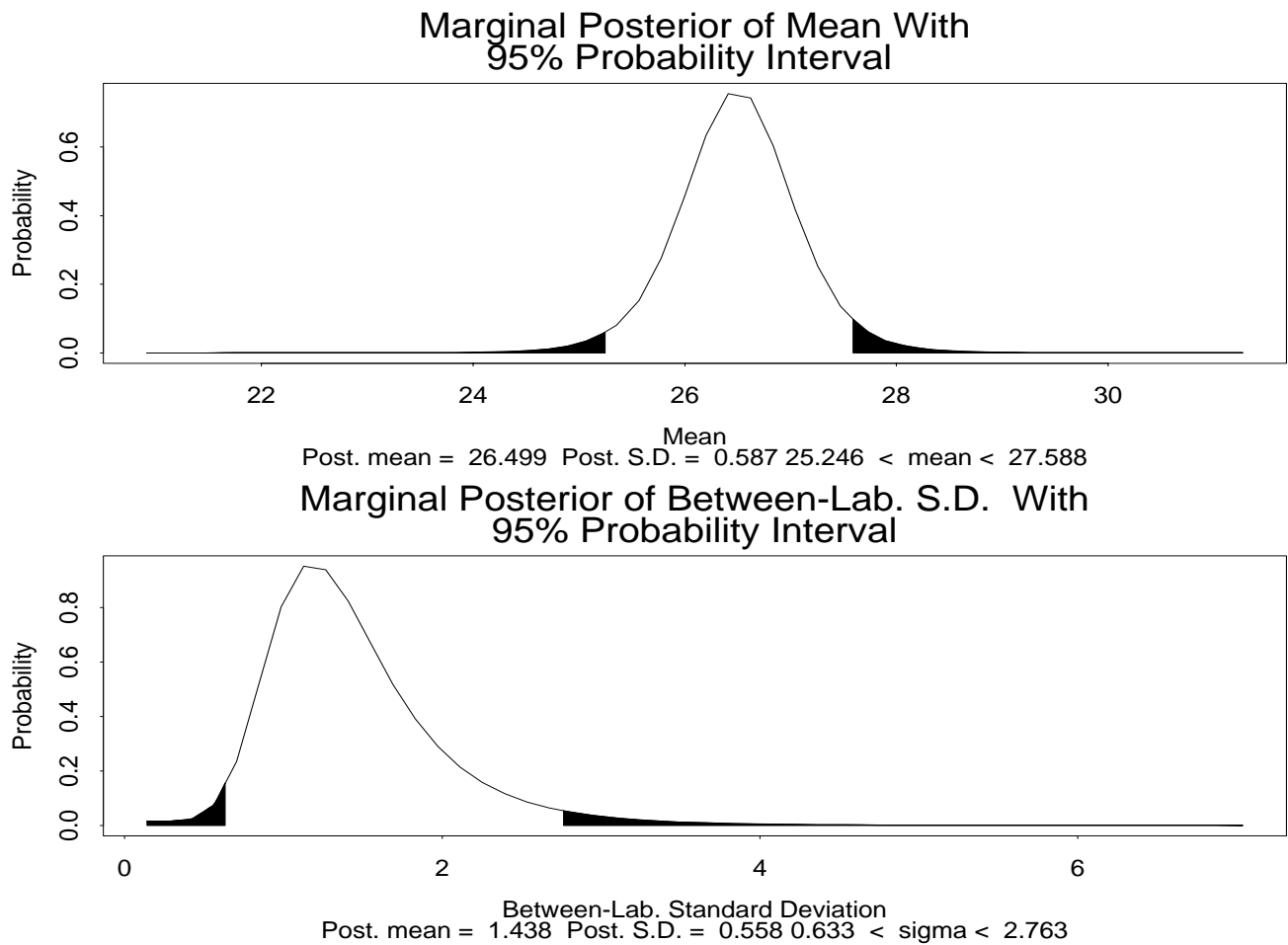
$$p(\mu, \sigma | \{x_{ij}\}) \propto p(\sigma) \prod_{i=1}^p \frac{1}{t_i} f_{n_i-1} \left[ \frac{x_i - \mu}{t_i}; \frac{2\sigma^2}{t_i^2} \right].$$

- The posterior of  $\mu$  given  $\sigma = 0$  is a product of scaled  $t$ -densities centered at the  $x_i$ , since

$$\frac{1}{t_i} f_{n_i-1} \left[ \frac{x_i - \mu}{t_i}; 0 \right] = \frac{1}{t_i} T'_{n_i-1} \left( \frac{x_i - \mu}{t_i} \right).$$

- We will take  $p(\sigma) = 1$ , though an arbitrary proper prior does not introduce additional difficulties.

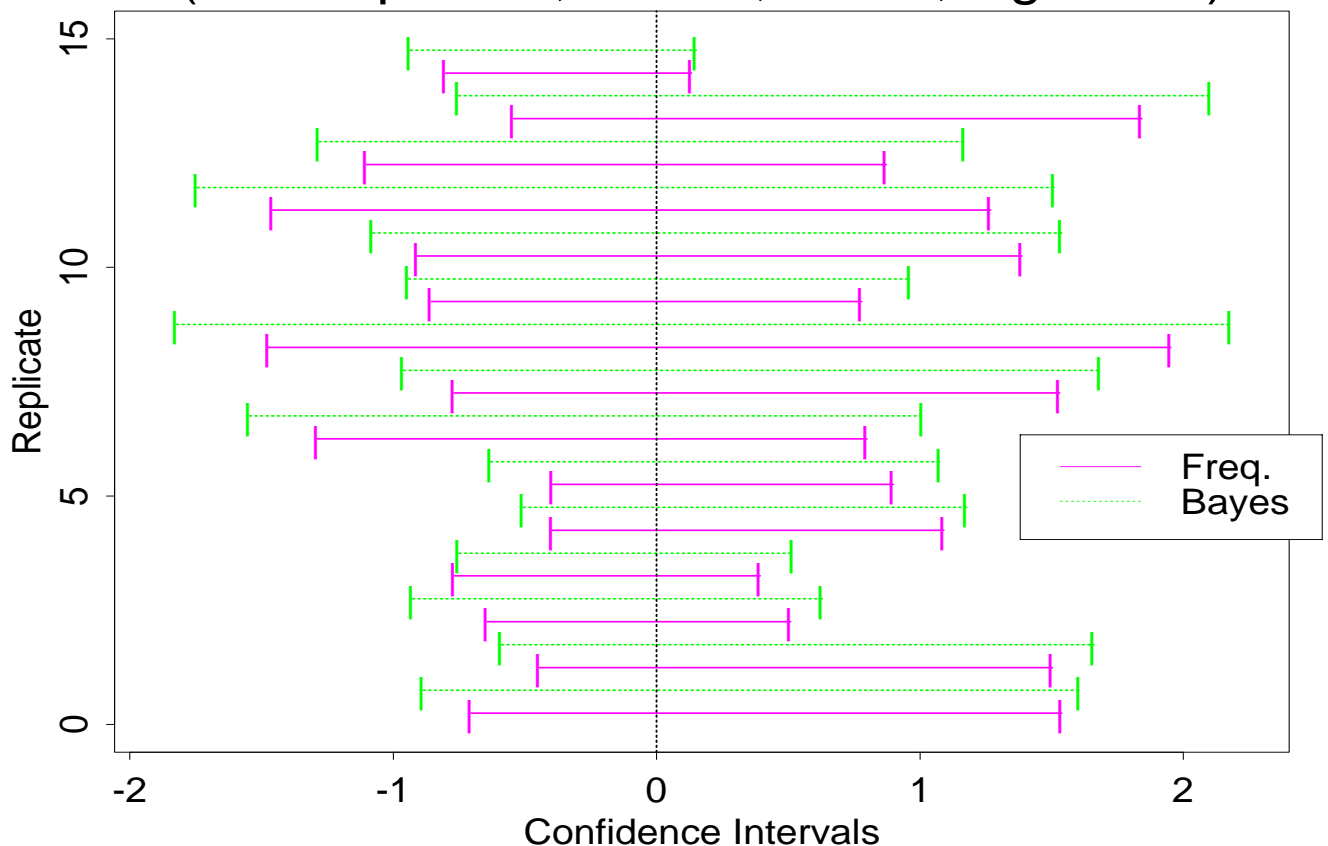
## Approximate Confidence Intervals: Apricot Fiber Data



## Small Simulation Comparing Bayesian and Frequentist Intervals

$$\begin{aligned}\mu &= 0 \\ \sigma_i &= \sigma_e \\ \sigma^2 + \sigma_e^2 &= 1 \\ \rho &= \sigma^2 / (\sigma_e^2 + \sigma^2) = 1/2\end{aligned}$$

Simulation Comparing Confidence Intervals  
(5 Groups of 5, rho=.5, mu=0, sigma =1)



## A Two-Way Mixed Model (Heteroscedastic, no Interaction)

$$x_{ijk} = \theta_k + \delta_i + e_{ijk},$$

- $i = 1, \dots, p$  Laboratories
- $j = 1, \dots, n_i$  Replicates
- $k = 1, \dots, m$  Materials

$$\delta_i \sim N(0, \sigma^2)$$

$$e_{ijk} \sim N(0, \sigma_i^2)$$

Some notation:  $\tau_i^2 \equiv \sigma_i^2 / (n_i m)$ ,  $\nu_i \equiv n_i m - 1$ .

## ML Equations

$$\theta_k - \bar{\theta} \equiv \phi_k = \frac{\sum_{i=1}^p (\bar{x}_{i \cdot k} - \bar{x}_{i \cdot \cdot}) / \tau_i^2}{\sum_{i=1}^p 1 / \tau_i^2}$$

$$\bar{\theta} = \frac{\sum_{i=1}^p \gamma_i \bar{x}_{i \cdot \cdot}}{\sum_{i=1}^p \gamma_i}$$

$$\sigma^2 = \frac{\sum_{i=1}^p \gamma_i \left[ (\bar{x}_{i \cdot \cdot} - \bar{\theta})^2 + \frac{\nu_i t_i^2}{1 - \gamma_i} \right]}{\sum_{i=1}^p n_i}$$

Where  $\tau_i^2 \equiv \sigma_i^2 / (n_i m)$ ,  $\nu_i \equiv m n_i - 1$ ,  
 $\gamma_i \equiv \sigma^2 / (\sigma^2 + \tau_i^2)$ , and

$$t_i^2 \equiv \frac{\sum_{j,k} (x_{ijk} - \bar{x}_{i \cdot k})^2 + n_i \sum_k (\bar{x}_{i \cdot k} - \bar{x}_{i \cdot \cdot} - \phi_k)^2}{\nu_i n_i m}$$

## ML Equations (Cont'd)

The weights  $\{\gamma_i\}_{i=1}^p$  are roots of the cubic equations

$$\begin{aligned} &\gamma_i^3 - (a_i + 2)\gamma_i^2 + \\ &[(n_i m + 1)a_i + \nu_i b_i + 1]\gamma_i - \\ &n_i a_i = 0 \end{aligned}$$

where

$$a_i \equiv \frac{\sigma^2}{(\bar{x}_{i..} - \bar{\theta})^2}$$

and

$$b_i \equiv \frac{t_i^2}{(\bar{x}_{i..} - \bar{\theta})^2}.$$

## An ML Iteration

1. Begin with estimates  $\left\{ \gamma_i^{(s)} \right\}$ .

2. Calculate the following:

$$\begin{aligned}\phi_k^{(s+1)} &= \frac{\sum_{i=1}^p (\bar{x}_{i.k} - \bar{x}_{i..}) / \tau_i^{2(s)}}{\sum_{i=1}^p 1 / \tau_i^{2(s)}} \\ \bar{\theta}^{(s+1)} &= \frac{\sum_{i=1}^p \gamma_i^{(s)} \bar{x}_{i..}}{\sum_{i=1}^p \gamma_i^{(s)}} \\ \sigma_{(s+1)}^2 &= \frac{\sum_{i=1}^p \gamma_i^{(s)} \left[ (\bar{x}_{i..} - \bar{\theta})^2 + \frac{\nu_i t_i^2}{1 - \gamma_i^{(s)}} \right]}{\sum_{i=1}^p n_i}\end{aligned}$$

3. Note that if the  $\phi_k$  are constrained to satisfy the above ML equation, then

$$t_i^2 = \frac{\sum_{j,k} (x_{ijk} - \bar{x}_{i..})^2 - \sum_k \phi_k^2 / m}{n_i \nu_i m}$$

4. Solve the cubics for new estimates  $\gamma_i^{(s+1)}$ , and iterate.

## Some Theoretical Results for Two-Way Mixed Model

The one-way results discussed earlier generalize:

- Monotone convergence
- All stationary values of likelihood in box in  $(\mu, \sigma, \sum_k \phi_k^2)$  space.
- Exactly one weight  $\gamma_i \in [0, 1]$ , unless  $i$ th lab an outlier and  $n_i$  small
- Variances cannot be negative at solution to likelihood equation.

## Summary

- A reparametrization of the likelihood in the one-way heteroscedastic model leads to new insights in likelihood and Bayesian analyses.
- A procedure of Mandel and Paule is equivalent to a modified REML estimator of the mean in an one-way heteroscedastic model.
- Many of these results carry over to two-way models; this work is ongoing.